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RAW SEQUENCE LISTING

DATE: 03/21/2003

PATENT APPLICATION: US/10/042,746

TIME: 14:30:58

Input Set : N:\Crif3\RULE60\10042746.RAW.txt

Output Set: N:\CRF4\03212003\J042746.raw

SEQUENCE LISTING

3 (1) GENERAL INFORMATION:

(i) APPLICANT: Drucker, Daniel J.

(ii) TITLE OF INVENTION: GLUCAGON-LIKE PEPTIDE 2 AND ITS THERAPEUTIC USE

(iii) NUMBER OF SEQUENCES: 4

(iv) CORRESPONDENCE ADDRESS:

(A) ADDRESSEE: Pennie & Edmonds LLP

(B) STREET: 1155 Avenue of the Americas

(C) CITY: New York

(D) STATE: New York

(E) COUNTRY: USA

(F) TEL: 1-212-361-1711

(v) COMPUTER READABLE FORM:

(A) MEDIUM TYPE: Floppy disk

(B) COMPUTER: IBM PC compatible

(C) OPERATING SYSTEM: PC-DOS/MS-DOS

(D) SOFTWARE: Patentix Release #1.0, Version #1.30

(vi) CURRENT APPLICATION DATA:

(A) APPLICATION NUMBER: US/10/042,746

(B) FILING DATE: 20-Nov-2002

(C) CLASSIFICATION: 514

(vii) PRIOR APPLICATION DATA:

(A) APPLICATION NUMBER: US/08/632,533

(B) FILING DATE: 12-APR-1996

(viii) ATTORNEY/AGENT INFORMATION:

(A) NAME: Friebe, Thomas E.

(B) REGISTRATION NUMBER: 29,258

(C) REFERENCE/WORKS NUMBER: 8607-004

(ix) TELECOMMUNICATION INFORMATION:

(A) TELEPHONE: (212) 790-9090

(B) TELEFAX: (212) 869-8864/9741

45 (2) INFORMATION FOR SEQ ID NO: 1:

(i) SEQUENCE CHARACTERISTICS:

(A) LENGTH: 37 amino acids

(B) TYPE: amino acid

(C) STRANDEDNESS:

(D) TOPOLOGY: unknown

(ii) MOLECULE TYPE: peptide

(ix) FEATURE:

(A) NAME/KEY: Modified-site

(B) LOCATION: 1

(D) OTHER INFORMATION: /note= "An amino acid selected from His,

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```

60 Arg, or Lys. Xaa may or may not be present in the sequence."
61 (ix) FEATURE:
62 (A) NAME/KEY: Modified-site
63 (B) LOCATION: 2
64 (D) OTHER INFORMATION: /note= "An amino acid selected from His,
65 Arg, or Lys. Xaa may or may not be present in the sequence."
66 (ix) FEATURE:
67 (A) NAME/KEY: Modified-site
68 (B) LOCATION: 15
69 (D) OTHER INFORMATION: /note= "An amino acid selected from Met,
70 Leu, Ile, Val or Cys."
71 (ix) FEATURE:
72 (A) NAME/KEY: Modified-site
73 (B) LOCATION: 17
74 (D) OTHER INFORMATION: /note= "An amino acid selected from Ala,
75 Ser, Thr, Pro, Gly, Asn, Asp, Glu or Gln."
76 (ix) FEATURE:
77 (A) NAME/KEY: Modified-site
78 (B) LOCATION: 18
79 (D) OTHER INFORMATION: /note= "An amino acid selected from Ala,
80 Ser, Thr, Pro or Gly."
81 (ix) FEATURE:
82 (A) NAME/KEY: Modified-site
83 (B) LOCATION: 19
84 (D) OTHER INFORMATION: /note= "An amino acid selected from His,
85 Arg or Lys."
86 (ix) FEATURE:
87 (A) NAME/KEY: Modified-site
88 (B) LOCATION: 19
89 (D) OTHER INFORMATION: /note= "An amino acid selected from Asn,
90 Asp, Glu, Gln, His, Arg or Lys."
91 (ix) FEATURE:
92 (A) NAME/KEY: Modified-site
93 (B) LOCATION: 16
94 (D) OTHER INFORMATION: /note= "An amino acid selected from His,
95 Arg or Lys. Xaa may or may not be present in the sequence."
96 (ix) FEATURE:
97 (A) NAME/KEY: Modified-site
98 (B) LOCATION: 27
99 (D) OTHER INFORMATION: /note= "An amino acid selected from His,
100 Arg or Lys. Xaa may or may not be present in the sequence."
101 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:
W--> 120 Xaa Xaa His Ala Asp Gly Ser Phe Ser Asp Glu Met Asn Thr Xaa Leu
121 1 5 10 15

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132     Asp Xaa Leu Ala Xaa Xaa Asp Phe Ile Asn Trp Leu Xaa Xaa Thr Lys
133           20           25           30
134     Ile Thr Asp Xaa Xaa
135           35
136 (2) INFORMATION FOR SEQ ID NO: 2:
137     (i) SEQUENCE CHARACTERISTICS:
138           (A) LENGTH: 37 amino acids
139           (B) TYPE: amino acid
140           (C) STRANDEDNESS:
141           (D) TOPOLOGY: unknown
142     (ii) MOLECULE TYPE: peptide
143     (ix) FEATURE:
144           (A) NAME/KEY: Modified site
145           (B) LOCATION: 1
146           (D) OTHER INFORMATION: /note= "An amino acid selected from His,
147 Arg, or Lys. Xaa may or may not be present in the sequence."
148     (ix) FEATURE:
149           (A) NAME/KEY: Modified-site
150           (B) LOCATION: 2
151           (D) OTHER INFORMATION: /note= "An amino acid selected from His,
152 Arg, or Lys. Xaa may or may not be present in the sequence."
153     (ix) FEATURE:
154           (A) NAME/KEY: Modified-site
155           (B) LOCATION: 11
156           (D) OTHER INFORMATION: /note= "An amino acid selected from Ala,
157 Ser, Thr, Pro or Gly."
158     (ix) FEATURE:
159           (A) NAME/KEY: Modified-site
160           (B) LOCATION: 36
161           (D) OTHER INFORMATION: /note= "An amino acid selected from His,
162 Arg, or Lys. Xaa may or may not be present in the sequence."
163     (ix) FEATURE:
164           (A) NAME/KEY: Modified-site
165           (B) LOCATION: 37
166           (D) OTHER INFORMATION: /note= An amino acid selected from His,
167 Arg, or Lys. Xaa may or may not be present in the sequence."
168 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:
W--> 173     Xaa Xaa His Ala Asp Gly Ser Phe Ser Asp Glu Met Asn Thr Ile Leu
174           1           10           15
175     Asp Asn Leu Ala Xaa Arg Asp Phe Ile Asn Trp Leu Ile Gln Thr Lys
176           16           25           30
177     Ile Thr Asp Xaa Xaa
178           31
179 (2) INFORMATION FOR SEQ ID NO: 3:
180     (i) SEQUENCE CHARACTERISTICS:
181           (A) LENGTH: 33 amino acids
182           (B) TYPE: amino acid
183           (C) STRANDEDNESS:
184           (D) TOPOLOGY: unknown

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189      (ii) MOLECULE TYPE: peptide
190      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:
196      His Ala Asp Gly Ser Phe Ser Asp Glu Met Asn Thr Ile Leu Asp Asn
197      1          5          10          15
198      Leu Ala Thr Arg Asp Phe Ile Asn Trp Leu Ile Gln Thr Lys Ile Thr
199      20          25          30
200      Asp
204 (2) INFORMATION FOR SEQ ID NO: 4:
206      (i) SEQUENCE CHARACTERISTICS:
207          (A) LENGTH: 33 amino acids
208          (B) TYPE: amino acid
209          (C) STRANDEDNESS:
210          (D) TOPOLOGY: unknown
211      (ii) MOLECULE TYPE: peptide
212      (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:
218      His Ala Asp Gly Ser Phe Ser Asp Glu Met Asn Thr Ile Leu Asp Asn
219      1          5          10          15
220      Leu Ala Ala Arg Asp Phe Ile Asn Trp Leu Ile Gln Thr Lys Ile Thr
221      20          25          30
222      Asp

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RAW SEQUENCE LISTING ERROR SUMMARY

DATE: 03/21/2003

PATENT APPLICATION: US/10/042,746

TIME: 14:30:59

Input Set : N:\Crf3\RULE60\10042746.RAW.txt

Output Set: N:\CRF4\03212003\J042746.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; Xaa Pos.1,2,15,18,21,22,29,30,36,37

Seq#:2; Xaa Pos.1,2,21,36,37

VERIFICATION SUMMARYPATENT APPLICATION: **US/10/042,746**

DATE: 03/21/2003

TIME: 14:30:59

Input Set : **N:\Crf3\RULE60\10042746.RAW.txt**Output Set: **N:\CRF4\03212003\J042746.raw**

L:28 M:220 C: Keyword misspelled or invalid format, [(A) APPLICATION NUMBER:]
L:29 M:220 C: Keyword misspelled or invalid format, [(B) FILING DATE:]
L:34 M:233 W: Alpha Fields not Ordered, Reordered [(C) CLASSIFICATION:] of (1)(vi)
L:120 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:0
M:341 Repeated in SeqNo=1
L:173 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:0
M:341 Repeated in SeqNo=2